

GenCore version 4.5  
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OM nucleic - nucleic search, using sw model  
Run on: April 5, 2000, 18:38:42 : Search time 6423.57 Seconds  
(without alignments)  
2513.359 Million cell updates/sec

Title: US-09-090-672B-1  
Perfect score: 4276  
Sequence: 1 TTTTACGGTTTTTCCCTCC.....ATCAGAAAAA 4276

Scoring table: OLIGO\_NUC  
Gapop 60.0 , Gapext 60.0

Searched: 4538634 seqs, 1887831982 residues

Word size : 0

Total number of hits satisfying chosen parameters: 9077268

Minimum DB seq length: 0  
Maximum DB seq length: 1000000

Post-processing: Listing first 45 summaries

Database : EST:\*

1:	em_est1:*
2:	em_est2:*
3:	em_est3:*
4:	em_est4:*
5:	em_est5:*
6:	em_est6:*
7:	em_est7:*
8:	em_est8:*
9:	em_est9:*
10:	em_est10:*
11:	em_est11:*
12:	em_est12:*
13:	em_est13:*
14:	em_est14:*
15:	em_est15:*
16:	em_est16:*
17:	em_est17:*
18:	em_est18:*
19:	em_est19:*
20:	gb_est1:*
21:	gb_est2:*
22:	gb_est3:*
23:	gb_est4:*
24:	gb_est5:*
25:	gb_est6:*
26:	gb_est7:*
27:	gb_est8:*
28:	gb_est9:*
29:	gb_est10:*
30:	gb_est11:*
31:	gb_est12:*
32:	gb_est13:*
33:	gb_est14:*
34:	gb_est15:*
35:	gb_est16:*
36:	gb_est17:*
37:	gb_est18:*
38:	gb_est19:*
39:	gb_est20:*
40:	gb_est21:*
41:	gb_est22:*
42:	gb_est23:*
43:	gb_est24:*

44:	gb_est25:*
45:	gb_est26:*
46:	gb_est27:*
47:	gb_est28:*
48:	gb_est29:*
49:	gb_est30:*
50:	gb_est31:*
51:	gb_est32:*
52:	em_est20:*
53:	em_est21:*
54:	em_est22:*
55:	em_est23:*
56:	em_est24:*
57:	em_est25:*
58:	em_est26:*
59:	gb_est33:*
60:	gb_est34:*
61:	gb_est35:*
62:	gb_est36:*
63:	gb_est37:*
64:	gb_est38:*
65:	em_est27:*
66:	em_est28:*
67:	em_est29:*
68:	em_est30:*
69:	gb_est39:*
70:	gb_est40:*
71:	gb_est41:*
72:	gb_est42:*
73:	gb_est43:*
74:	gb_est44:*
75:	em_est31:*
76:	em_est32:*
77:	em_est33:*
78:	em_est34:*
79:	gb_gss1:*
80:	gb_gss2:*
81:	gb_gss3:*
82:	gb_gss4:*
83:	em_gss1:*
84:	em_gss2:*
85:	em_gss3:*
86:	em_gss4:*
87:	gb_gss5:*
88:	gb_gss6:*
89:	gb_gss7:*
90:	gb_gss8:*
91:	gb_gss9:*
92:	em_gss5:*
93:	em_gss6:*
94:	em_gss7:*
95:	em_gss8:*
96:	em_gss9:*
97:	em_gss10:*
98:	em_gss11:*
99:	gb_gss10:*
100:	gb_gss11:*
101:	em_gss12:*
102:	gb_gss12:*
103:	gb_gss13:*
104:	gb_gss14:*
105:	gb_gss15:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Score	Match Length	ID	Description
C 1	386	9.0	506 45	AI392674 t947c02.x